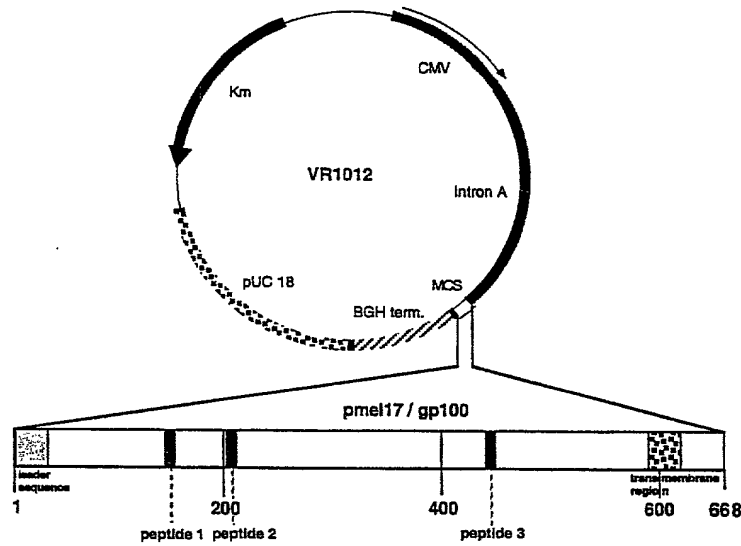


A



B

human	1	MDLVLRRCILRRAVILGALLAVGATKVPNRQDWLGVSRLRTRKAWNRQLYPEWTEACRLDC
mouse	1	MVGVRRRSELPVILVESAIIAVGALEGSRNQDWLGVPRLVTRKAWNRQLYPEWTEVCGSNC
human	61	WRGGOVSLKVINNDGPTLIGANASFSTALNFFPGSQKVLDPGQVIVNNTTIINGSQVWGGQP
mouse	61	WRGGOVSLKVINNDGPTLIGANASFSTALNFFPGSQKVLDPGQVIVNNTTIINGSQVWGGQP
human	121	VYPQETDDACHFPDGGPCPSCSWSCKRSEFVYVWKTWQYQVVLGGPVSGLSTIEHCHAMLC
mouse	121	VYPQETDDACHFPDGGPCPSCFKPKRKRSEFVYVWKTWQYQVVLGGPVSRSIATRHAKLG
human	181	THTMEVTVYHRRGSRSYVPLAHSSSAFTITDQVFFSVSVSOLRALDGCNKHFLRNQPTTF
mouse	181	THTMEVTVYHRRGSRSYVPLAHASTFTITDQVFFSVSVSOLCALDGETKHFLRNHPTTF
human	241	ALQLHDPSCGYLAEDLSYTWDFGDSGCTLLISRAFPVTHTYLEPCFVTAQVVLQAAIPLVS
mouse	241	ALQLHDPSCGYLAEDLSYTWDFGDCGCTLLISRALDVHTYLESFSVTAQVVLQAAIPLVS
human	301	CGSSPVPCTTDCGRPTAEAPNTTACQVPTTEVVGTTPCQAPTAEPSCTTSVQVPTTEVTS
mouse	301	CGSSPVPCTTDCYMPTEAPAGCTTSRQCTTKRVVGTTPCQMPPTOPSGTIVVQMPTEVTA
human	361	TAQVQMPTEASTGMPTEKVPVSQVLCITLAEMSTPEATGCTAEVSIIVLSCTTAQVTT
mouse	361	HTSQMLT-----SAVLDITLAEMSTPEATGCTPETRF-----SGTIVQATF
human	421	TEWVETTARELPPIPEPGPDASSINSTESTGSGCEPLIDGATTRLVKRQVPLDCVLYRY
mouse	403	TE-----CPDASPTPTPOSSFGSHSPLIDDQDTMLVKRQVPLDCVLYRY
human	481	GSFSTALDIVQGIESAEILQAVPFSCEGDAFELTVSCQGGLPKEACMETISSPGCQPPAQRIL
mouse	448	GSFSTALDIVQGIESAEILQAVPFSCEGDAFELTVSCQGGLPKEACMDISSPGCQPPAQRIL
human	541	QCPVLPSPACQVLVHQTLKGGSGTYCLNVSLADANSLAVVSTQLIMPVPGTLLTGOEAGL
mouse	508	QCSVPSPSPACQVLVHQTLKGGSGTYCLNVSLADANSLAVASTQLVWP-----GQGGGL
human	601	QGVPLIVGILLVIMAVVLASLIY-----RRRLMKQDFSYQPPHSSSHWLRLPRIFCSQPT
mouse	561	QCAPLIVGILLVIVAVVLASLILGIDLSRAOFFPKCHVAFTAAPASGHRA-R-----G-G
human	657	GENSPILLSQQQV
mouse	615	GENSPILLSQQQV

Fig. 1

A prophylactic 2/5

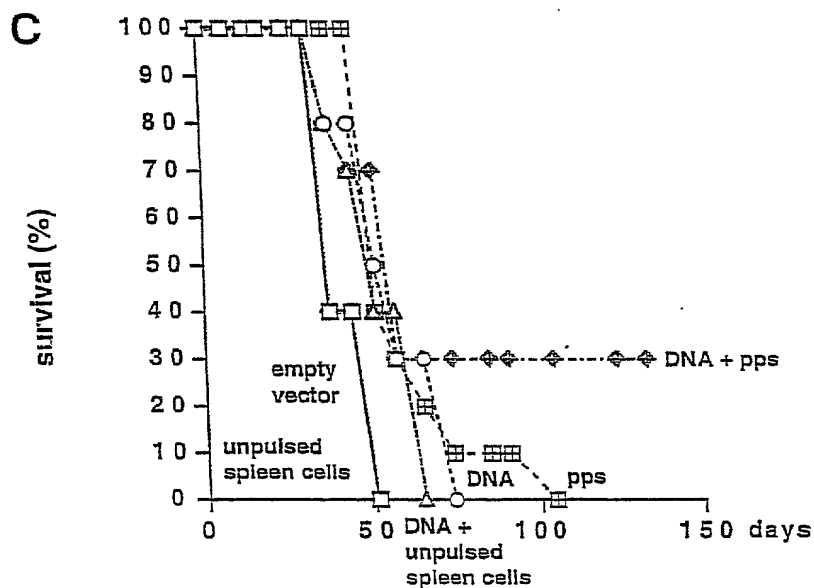
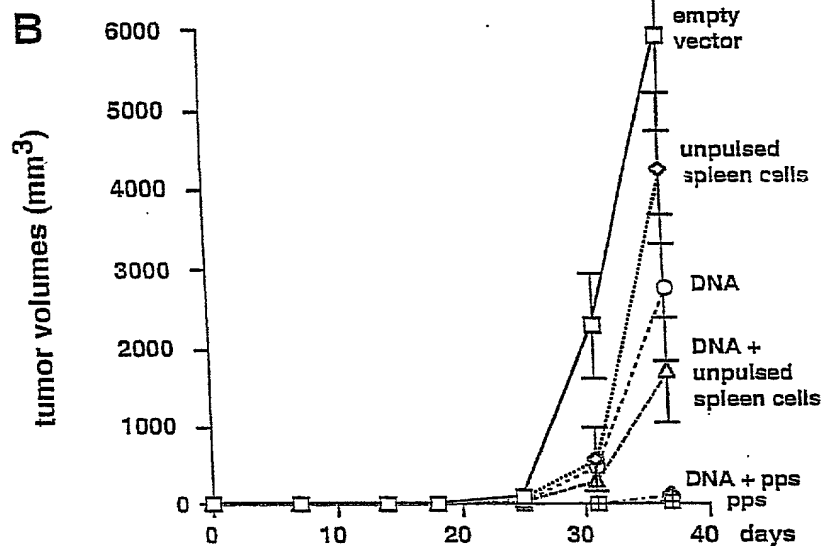
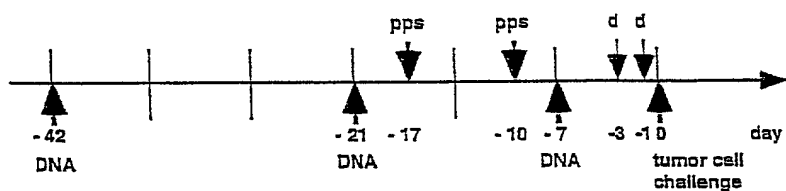


Fig. 2

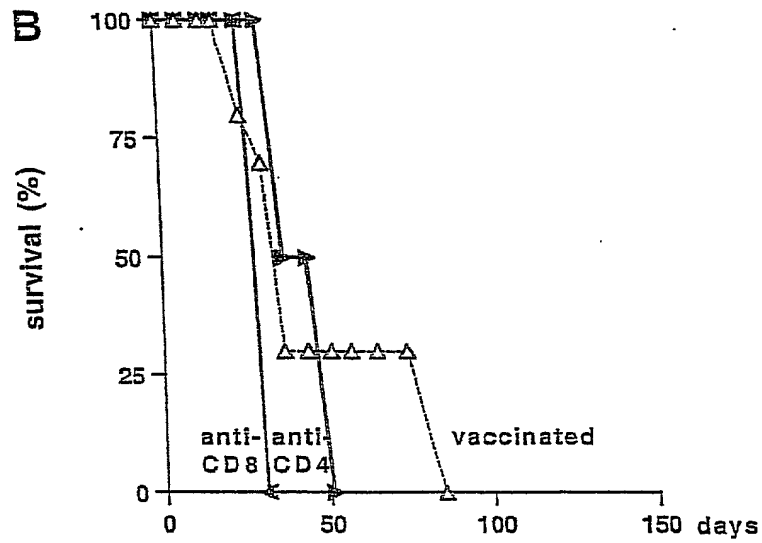
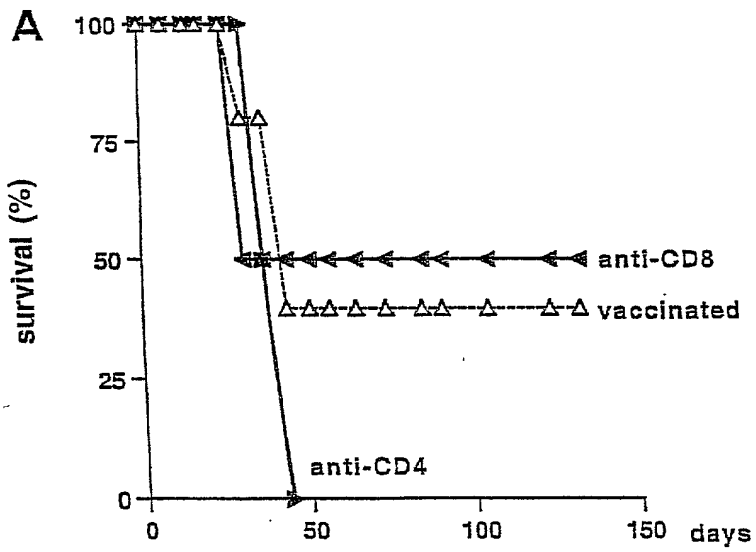


Fig. 3

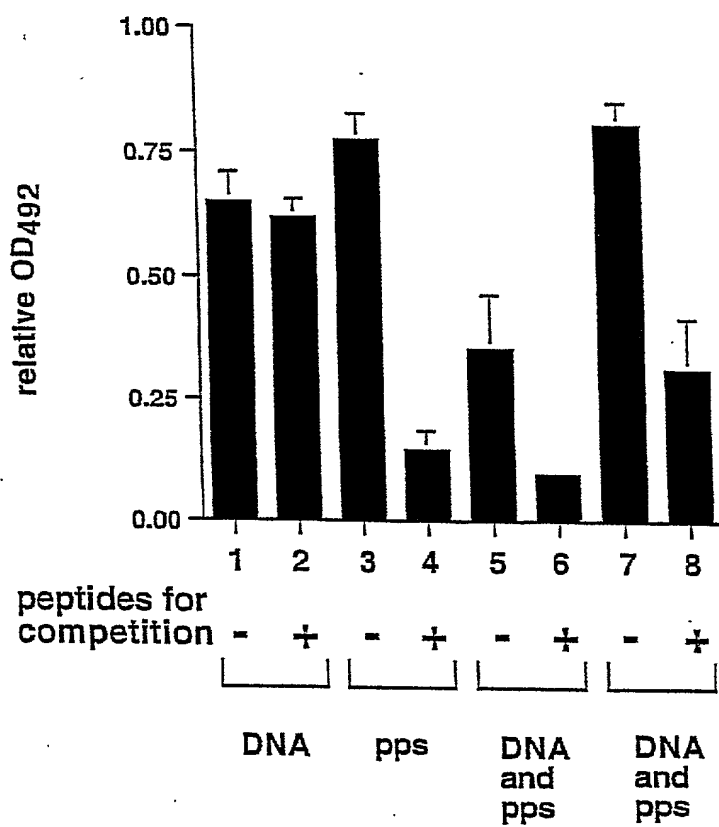


Fig. 4

A therapeutic 5/5

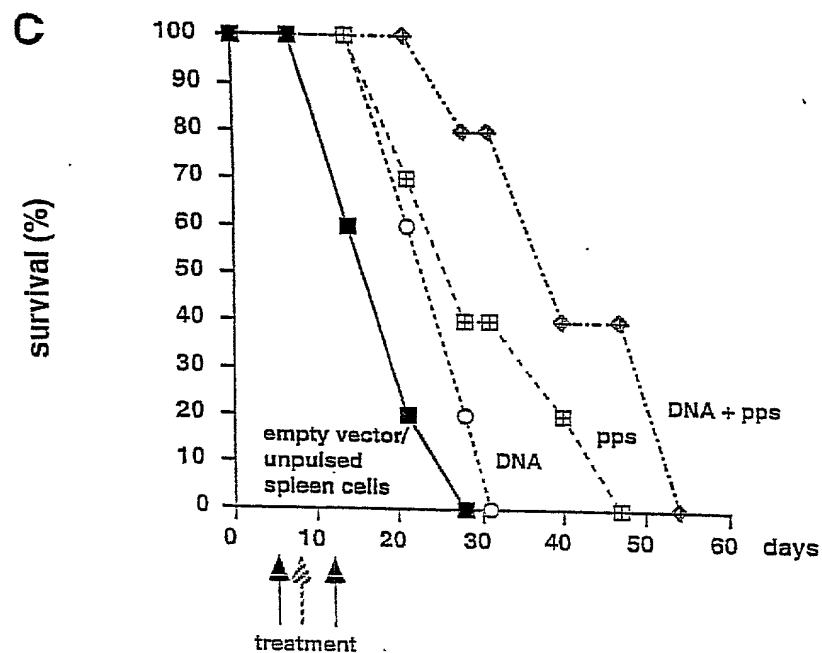
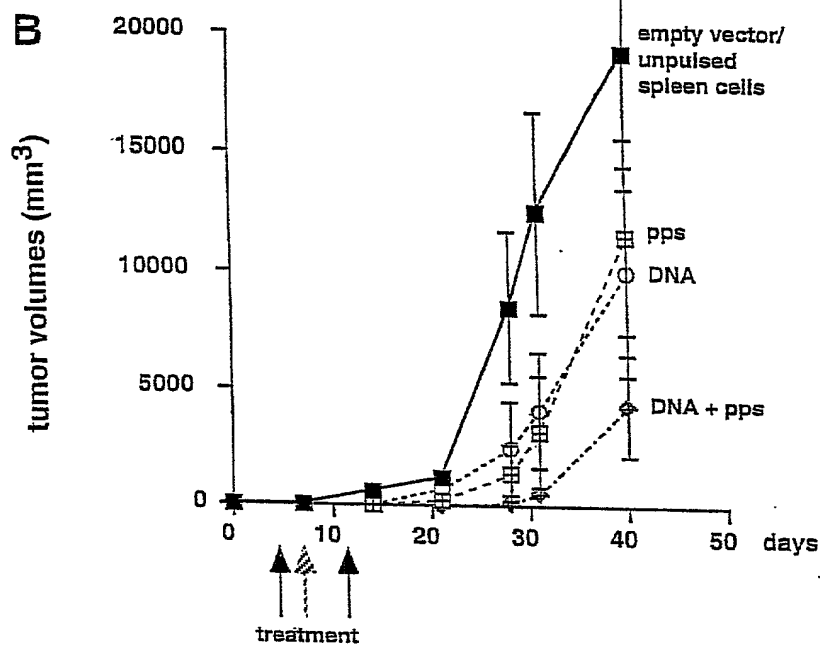
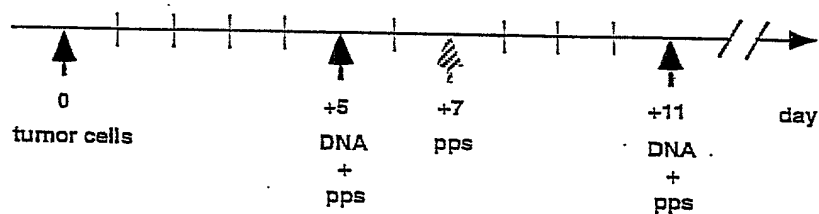


Fig. 5